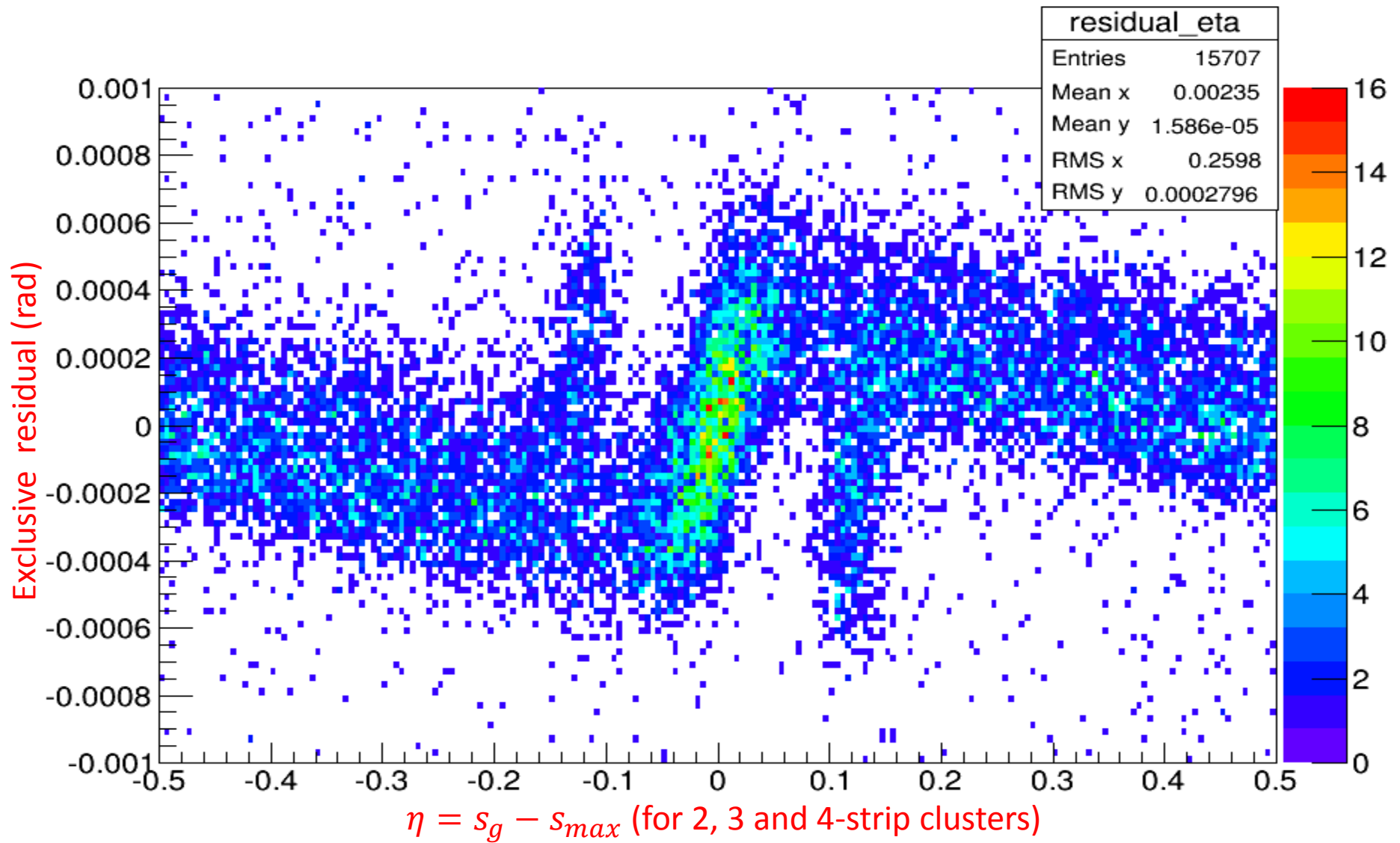


Position correction / resolution -- update

Aiwu Zhang, Marcus Hohlmann

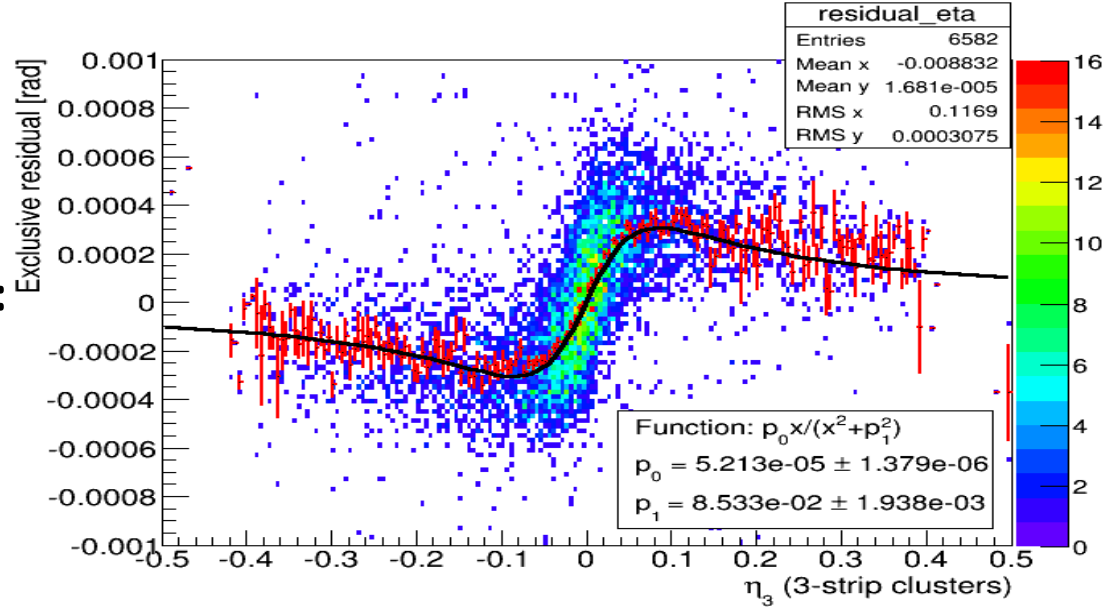
2014-08-18

HV scan data combined, 2,3,4-strip clusters are selected.

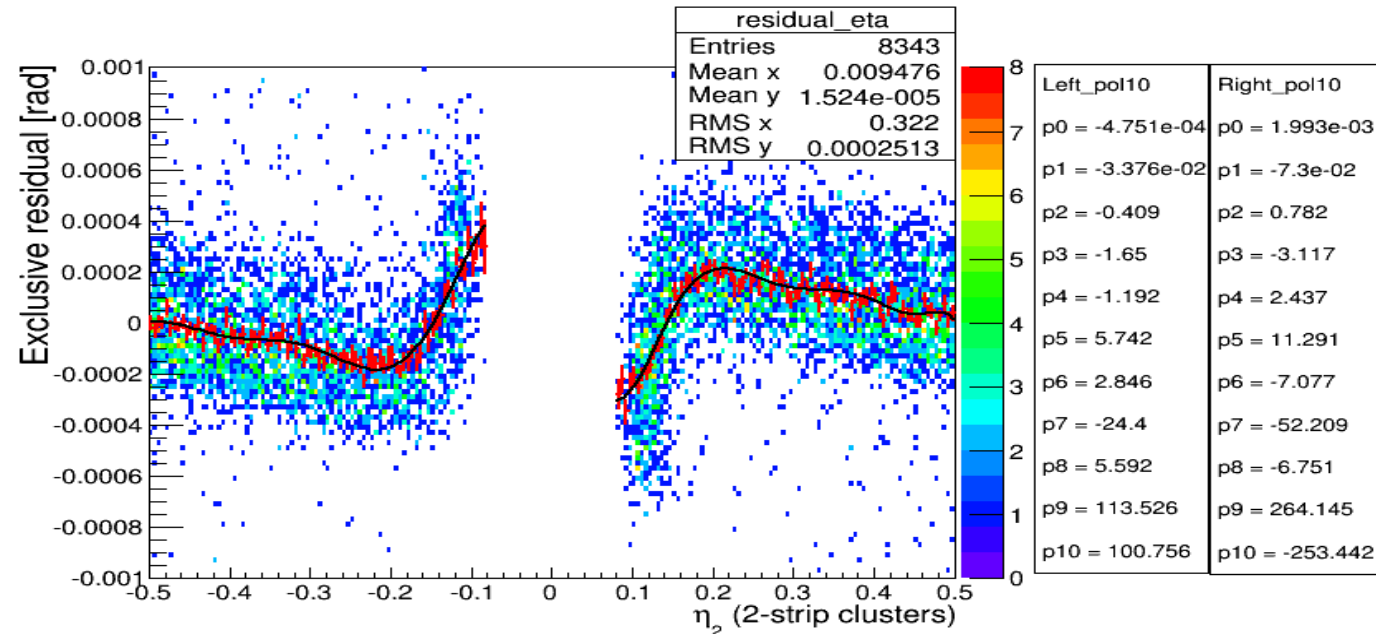


- Combined zigzag HV scan data, 90k raw events.
- Single cluster in every event in every detector; cluster size in trackers ≥ 1 .

3-strip clusters case:
Profile is fitted with
a **serpentine**
function

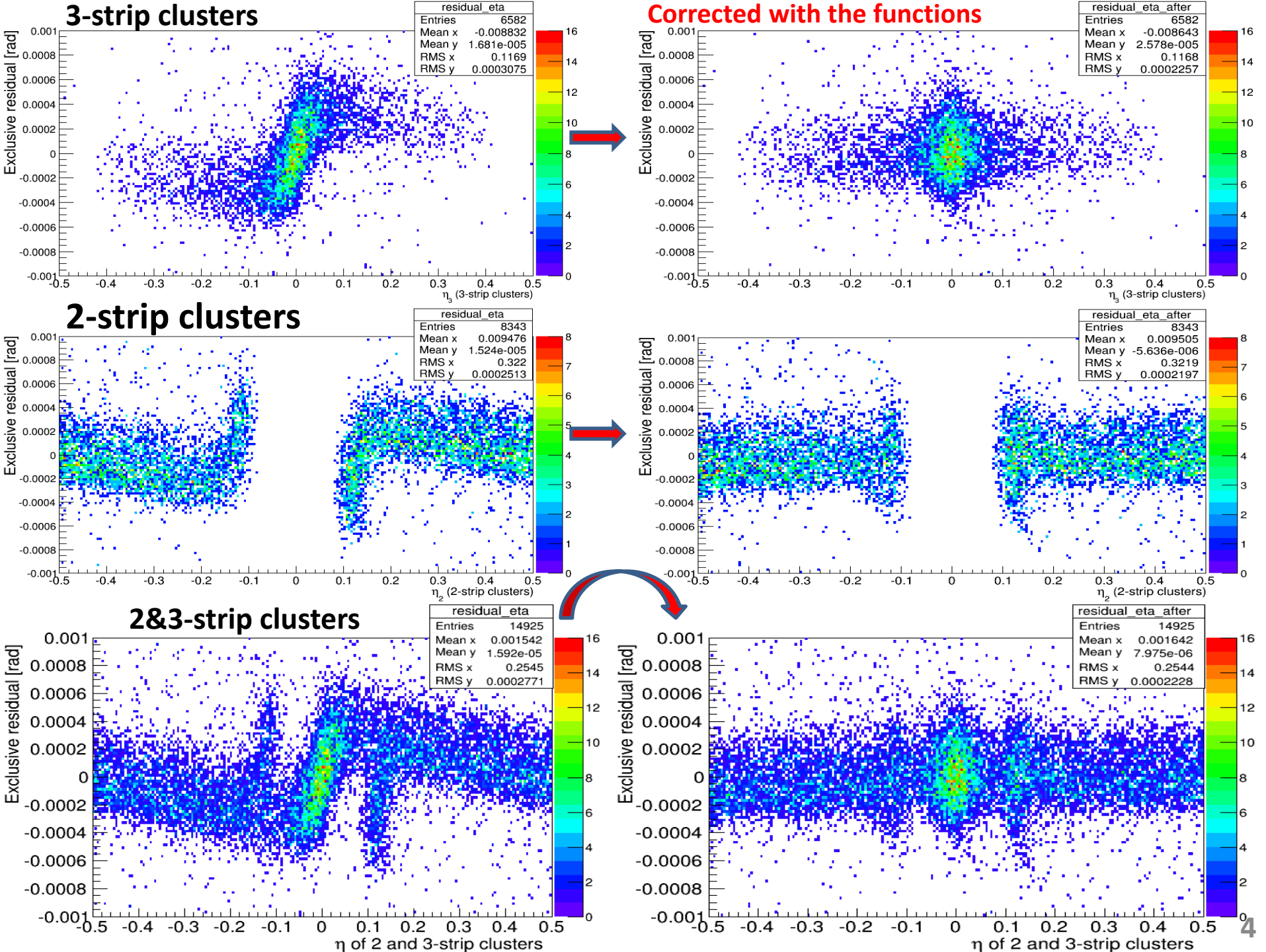


2-strip clusters case:
Profile is fitted with
two deci-**polynomials**
(degree 10)

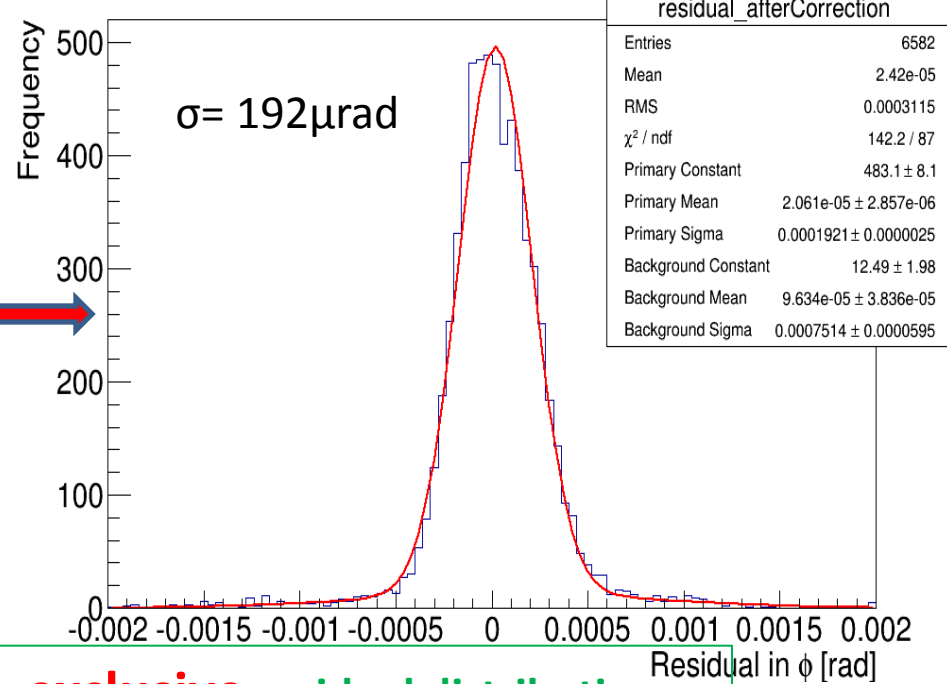
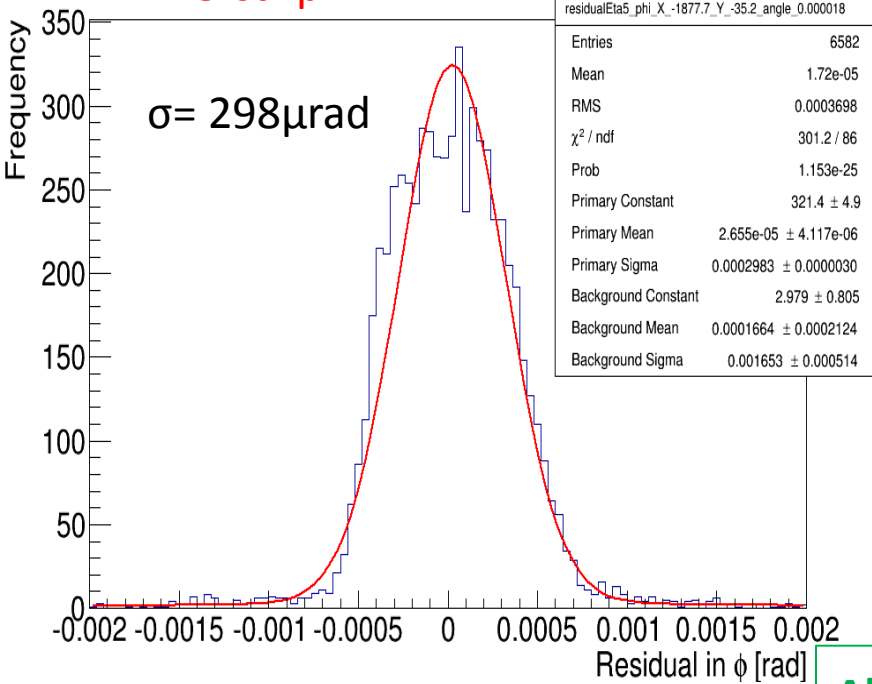


Residual = $\varphi_p - \varphi_m$, φ_p is the cluster position on the zigzag GEM given by trackers, φ_m is the position measured by the zigzag GEM.

Residual' = $\varphi_p - \varphi_m - f(\eta)$, pulls residual mean towards zero fall all η values.

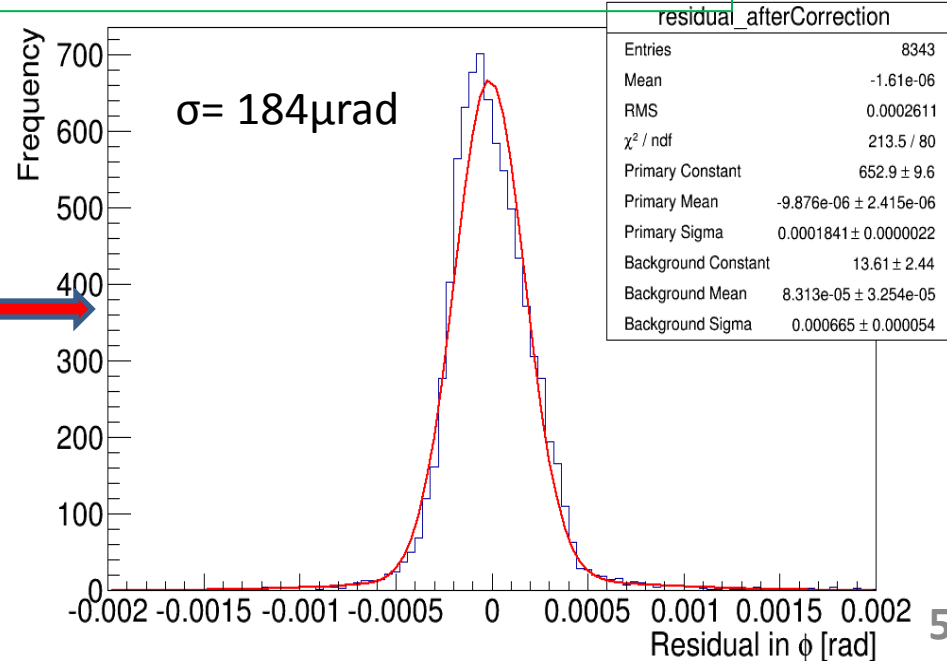
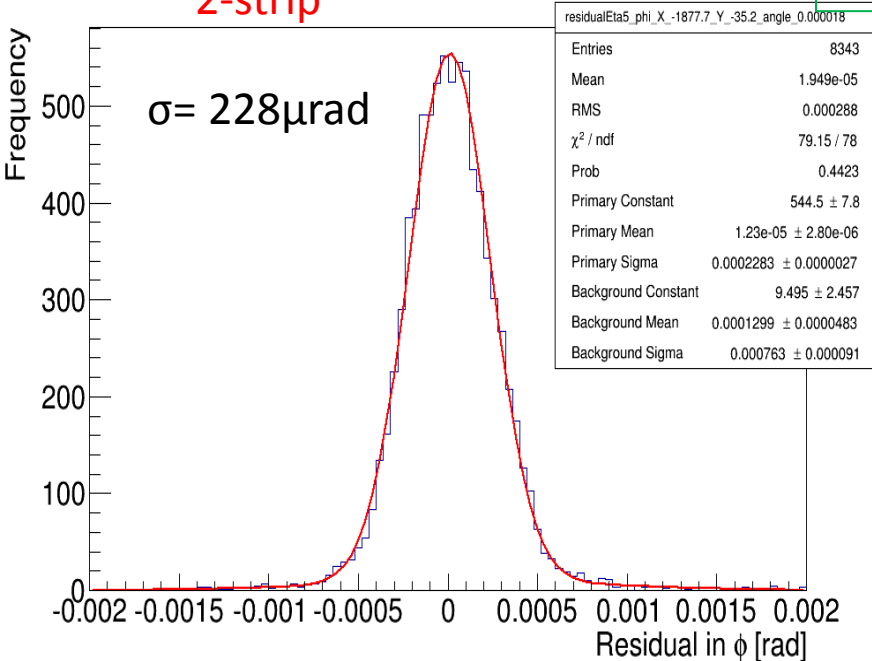


3-strip

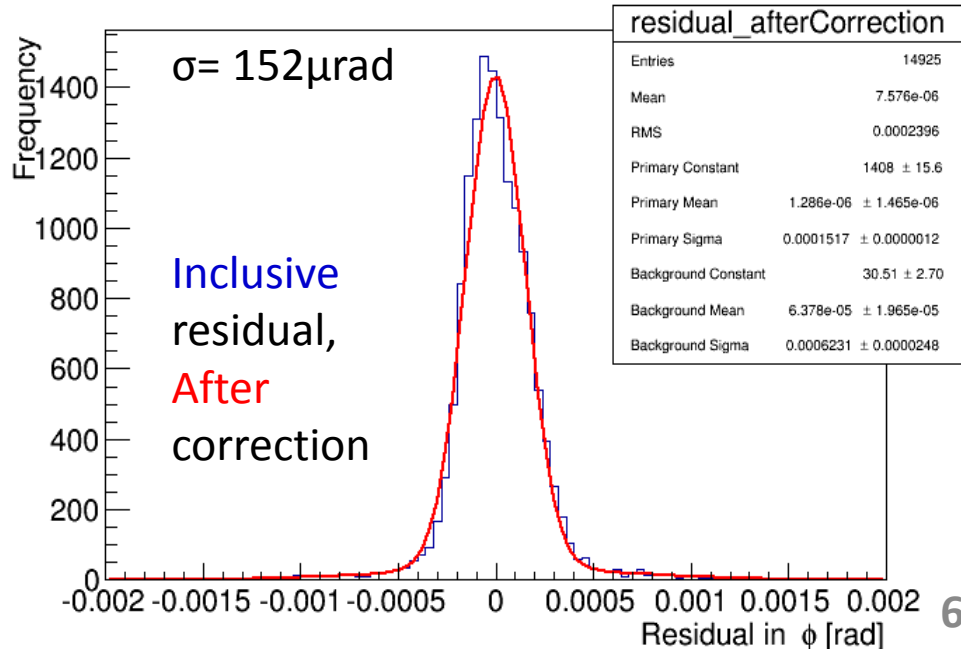
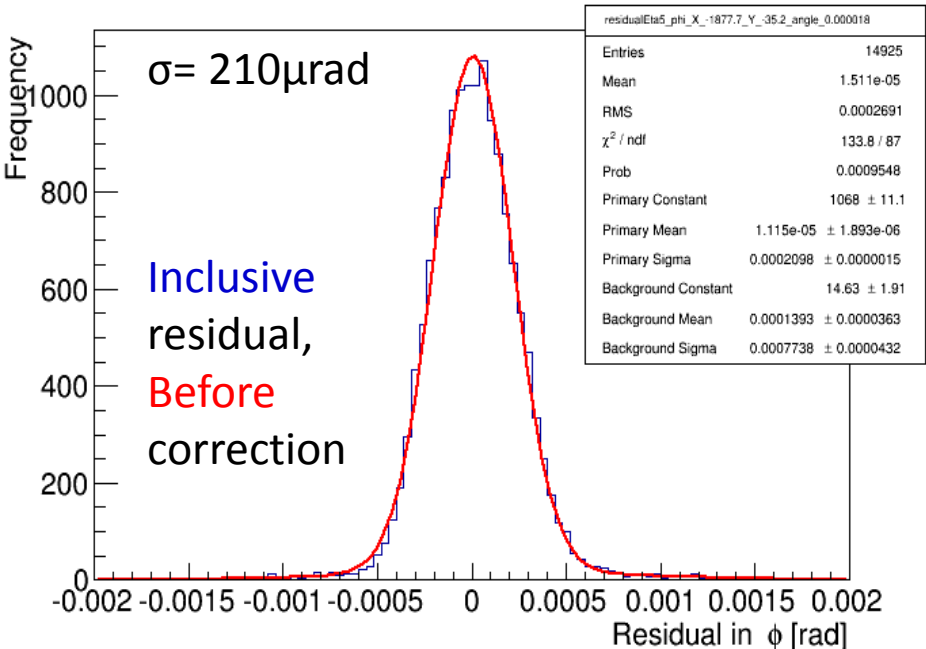
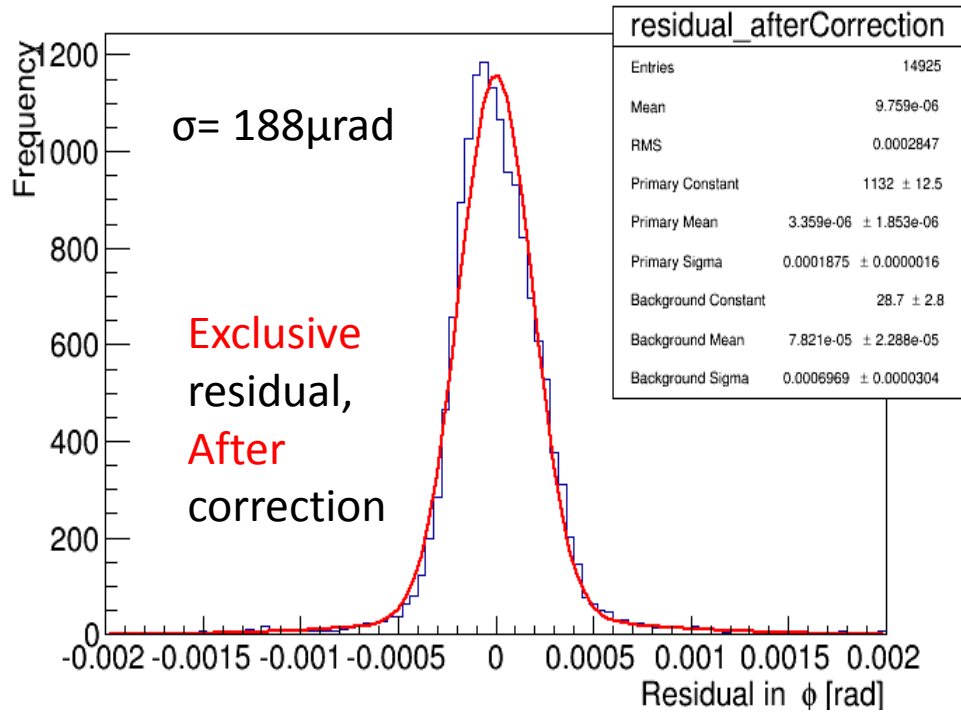
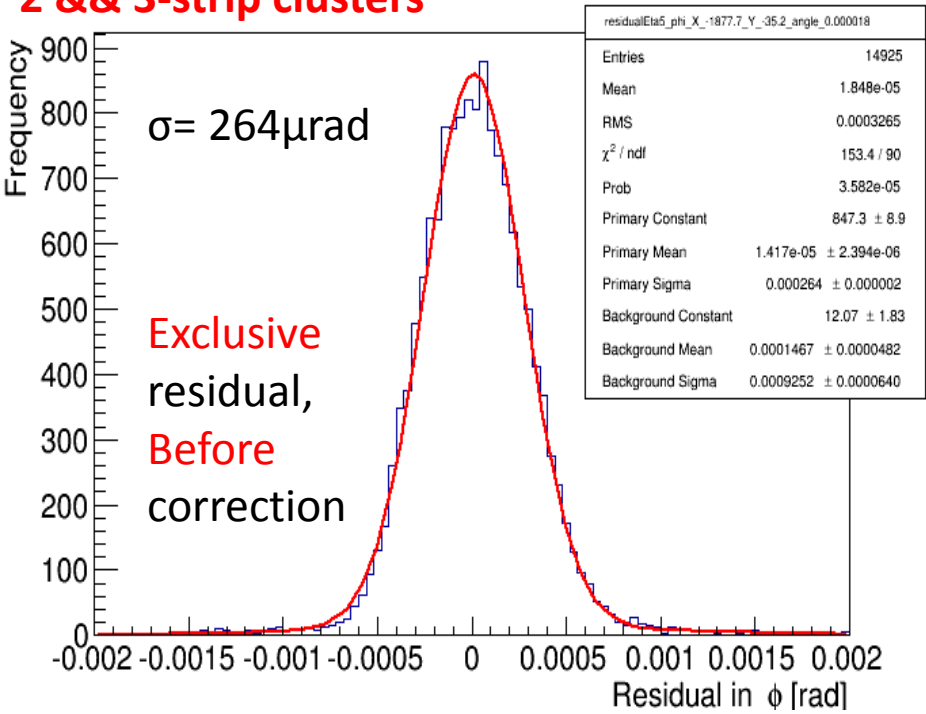


All are **exclusive** residual distributions

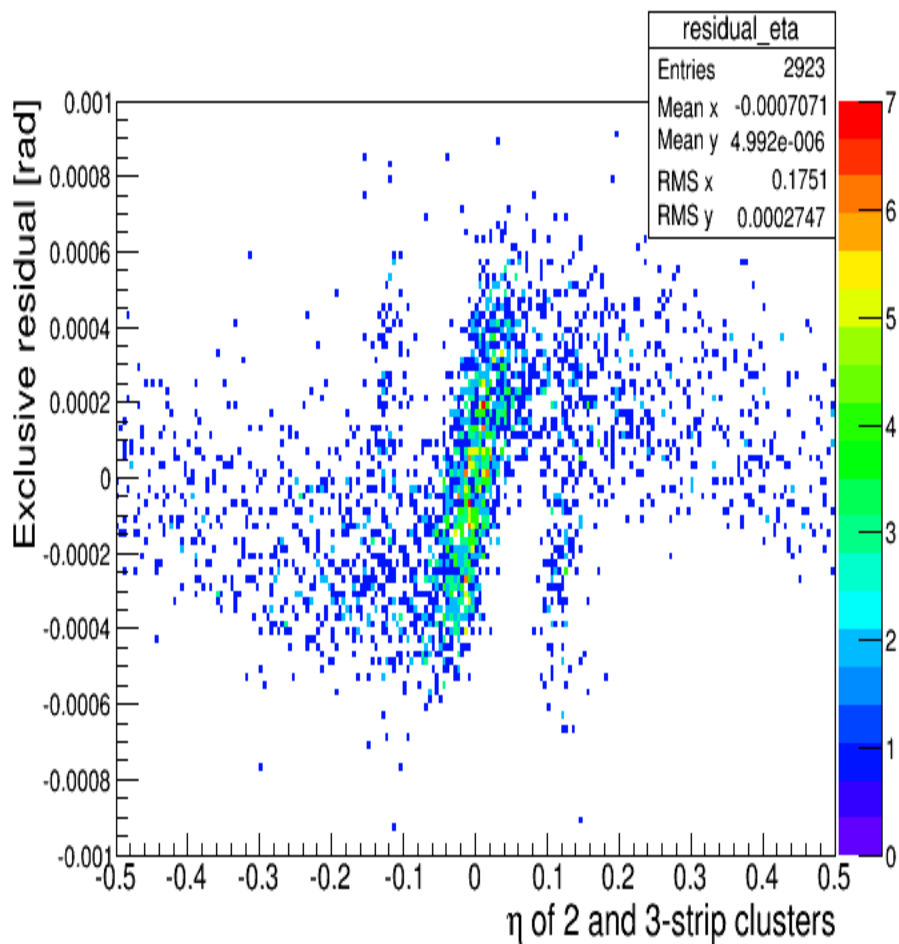
2-strip



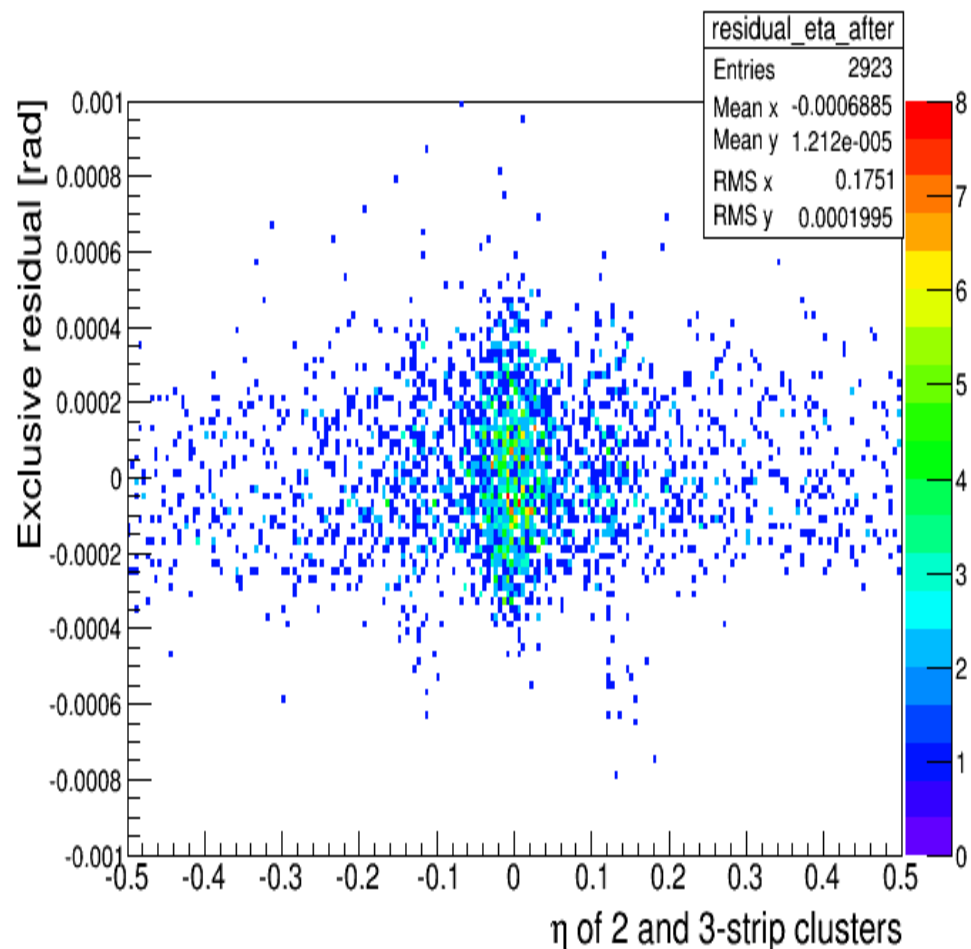
2 && 3-strip clusters



(exclusive) residual vs. eta plot @ 3400V for the zigzag (with 2 and 3-strip clusters)

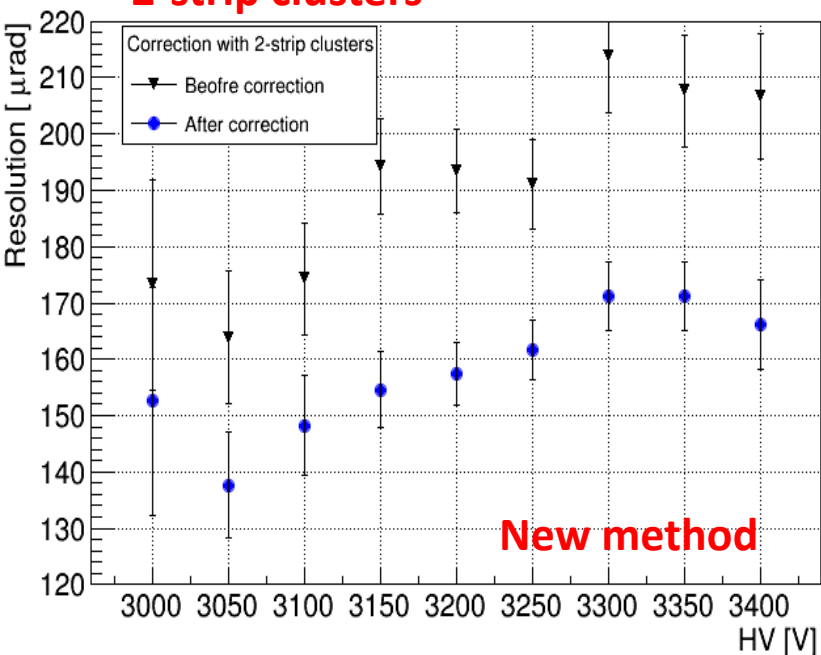


Before correction

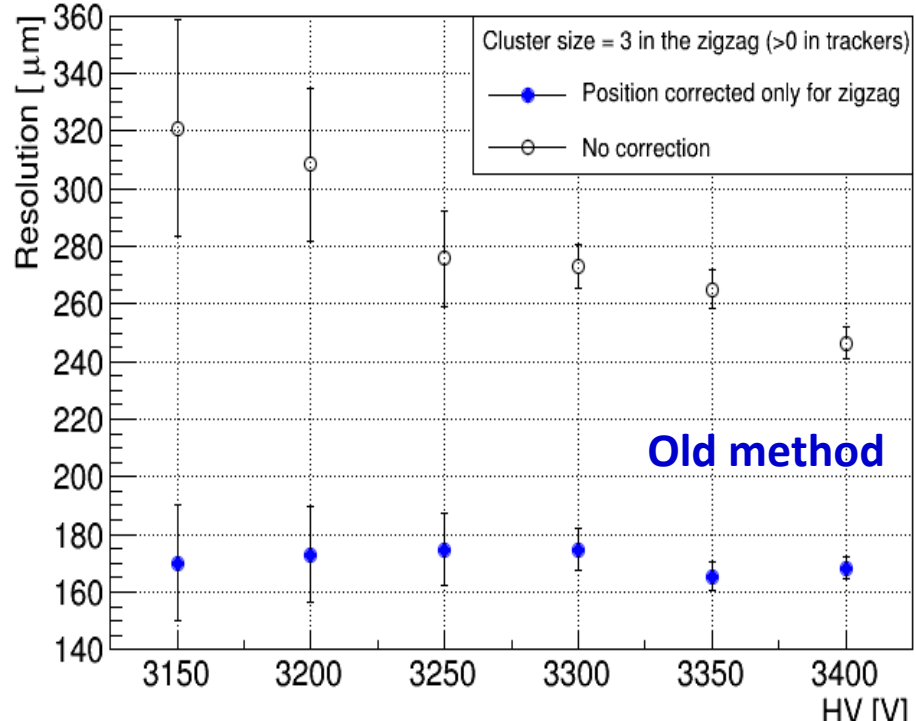
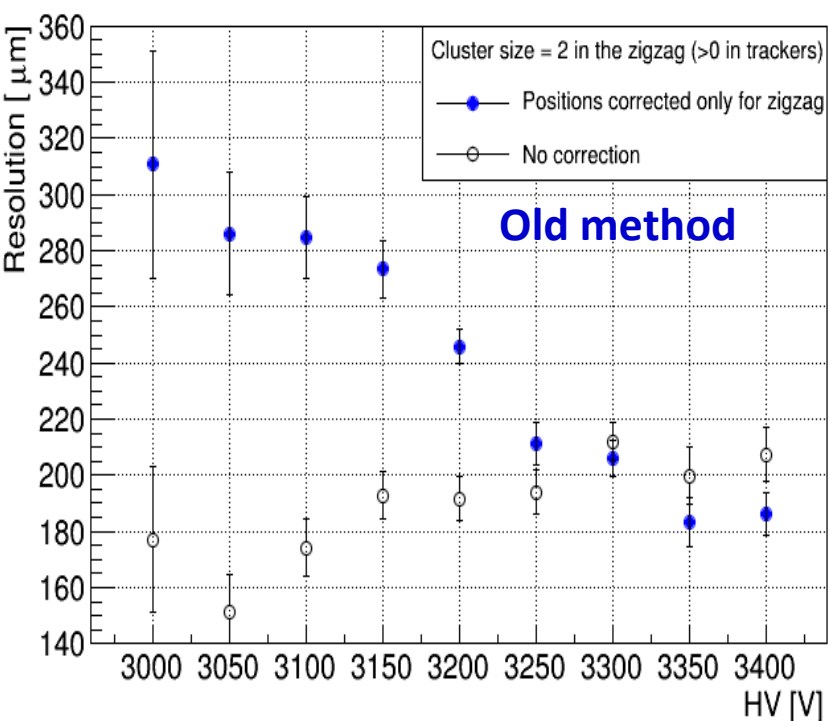
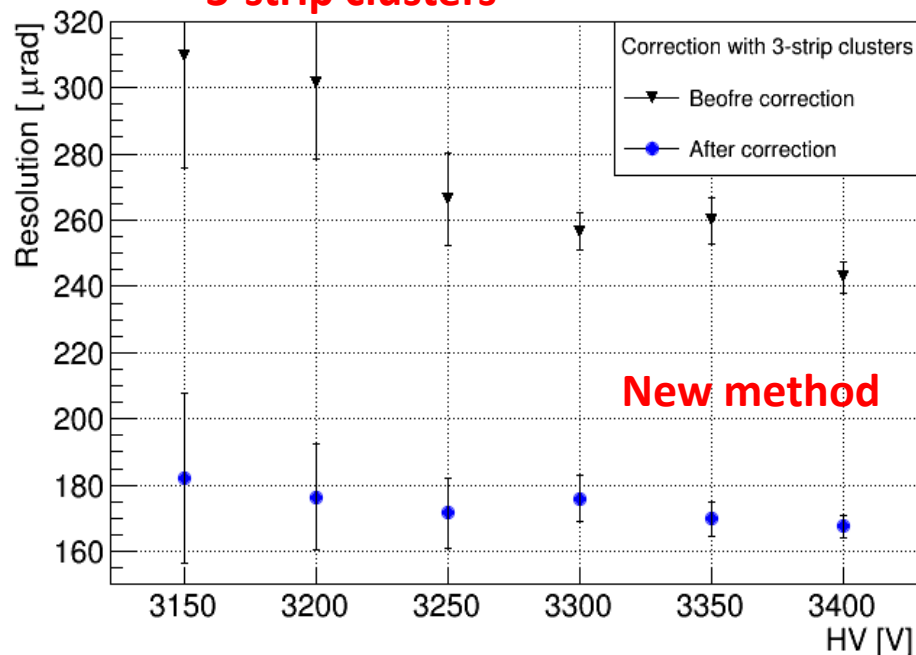


After correction

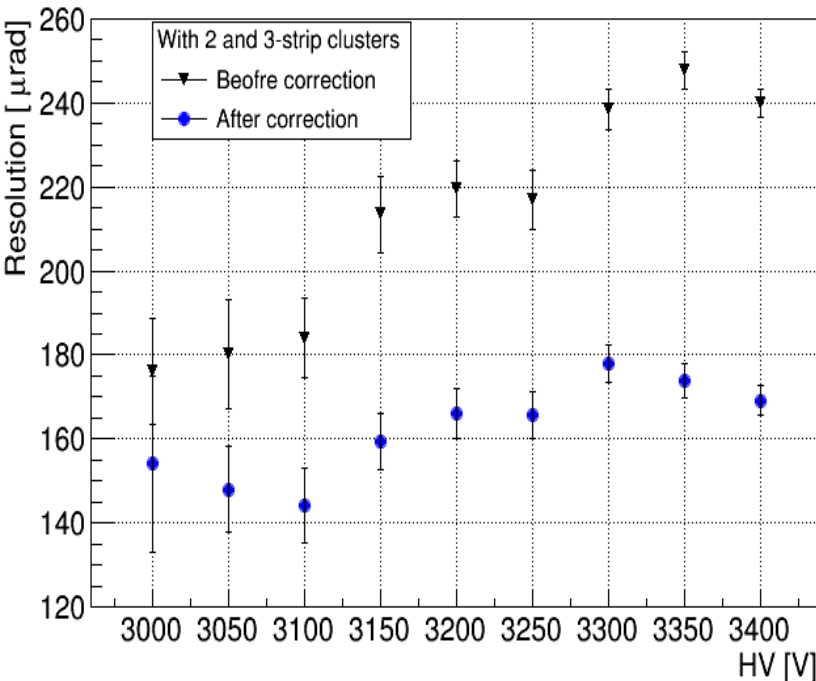
2-strip clusters



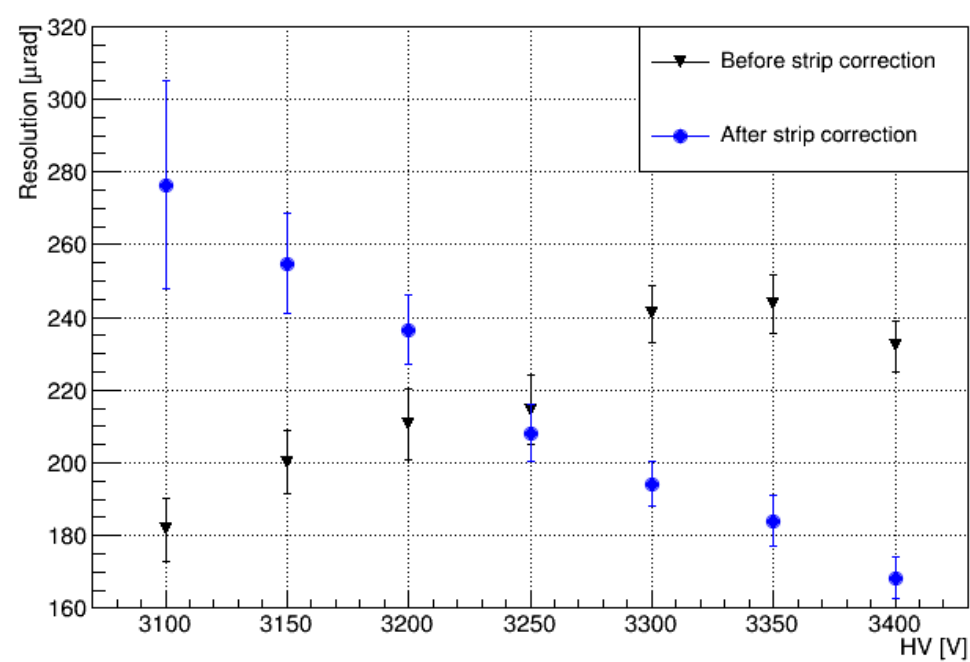
3-strip clusters



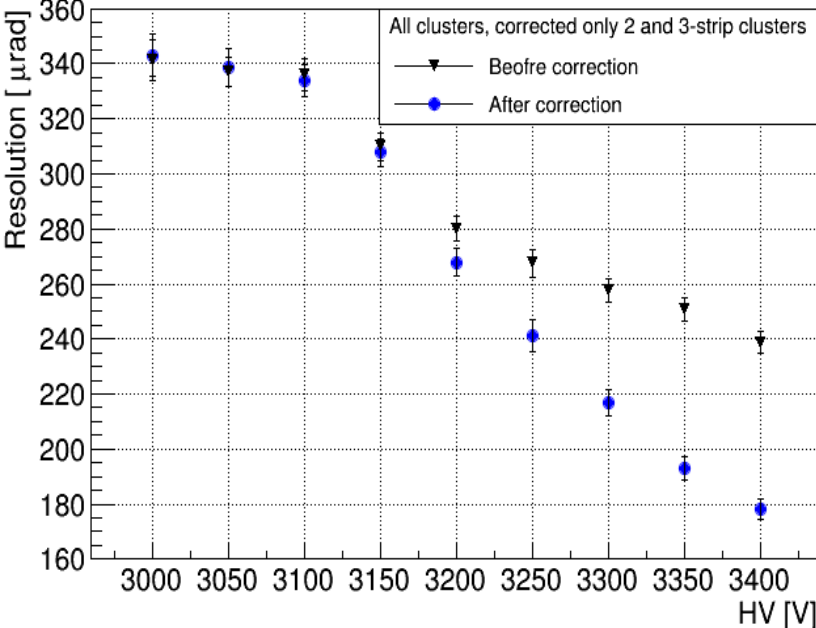
2 & 3-strip clusters **New method**



2, 3 & 4-strip clusters **Old method**

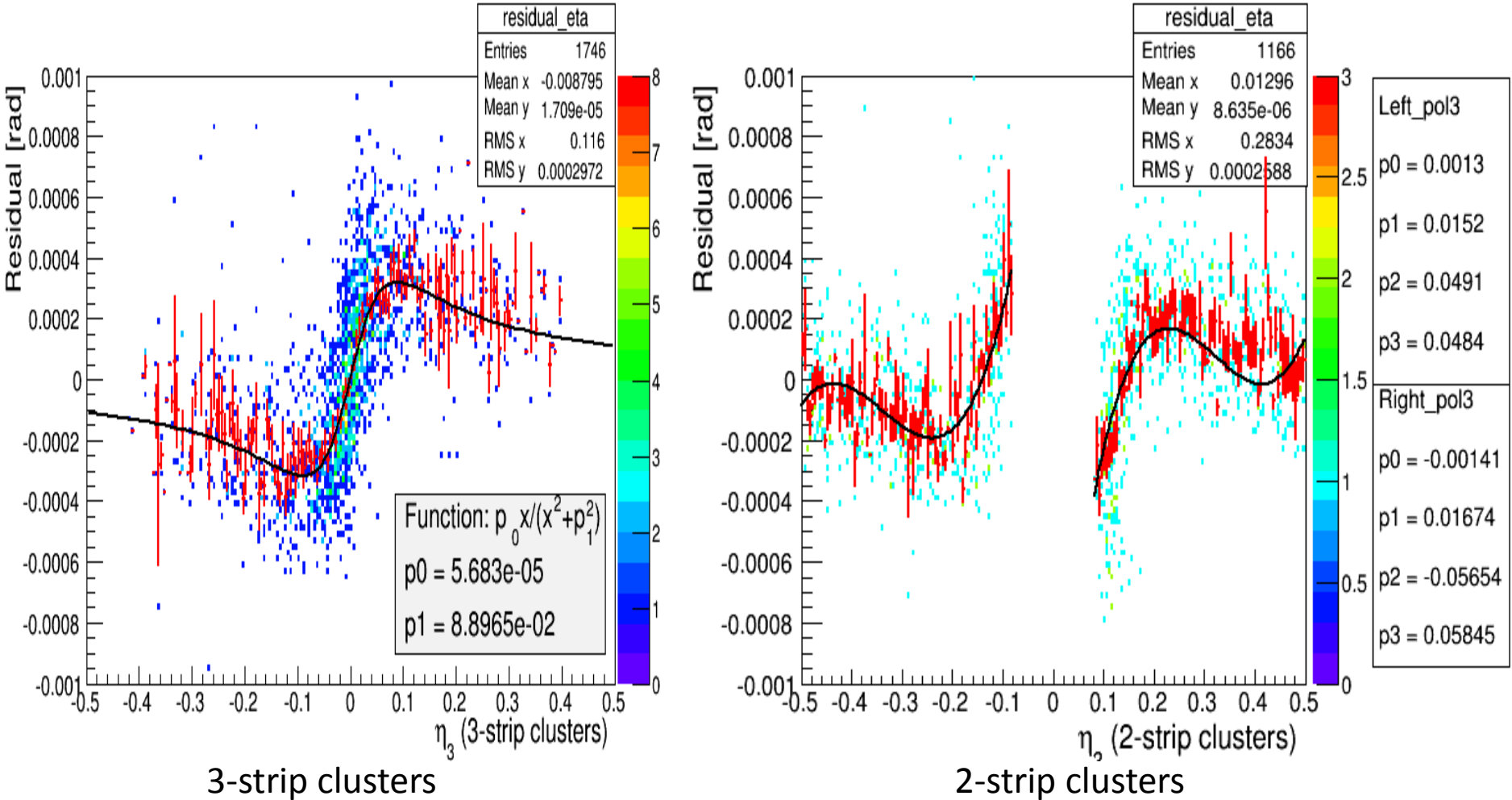


All clusters **New method**



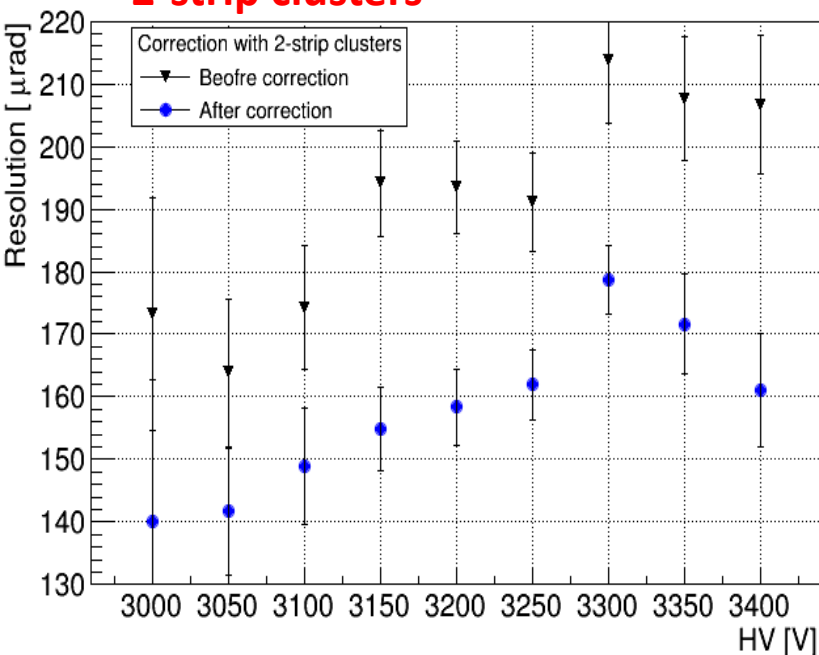
The plot on the left shows resolutions with all cluster sizes ($N \geq 1$):

No correction is applied for the black points;
Correction is (only) applied to 2 and 3-strip clusters for the blue points.

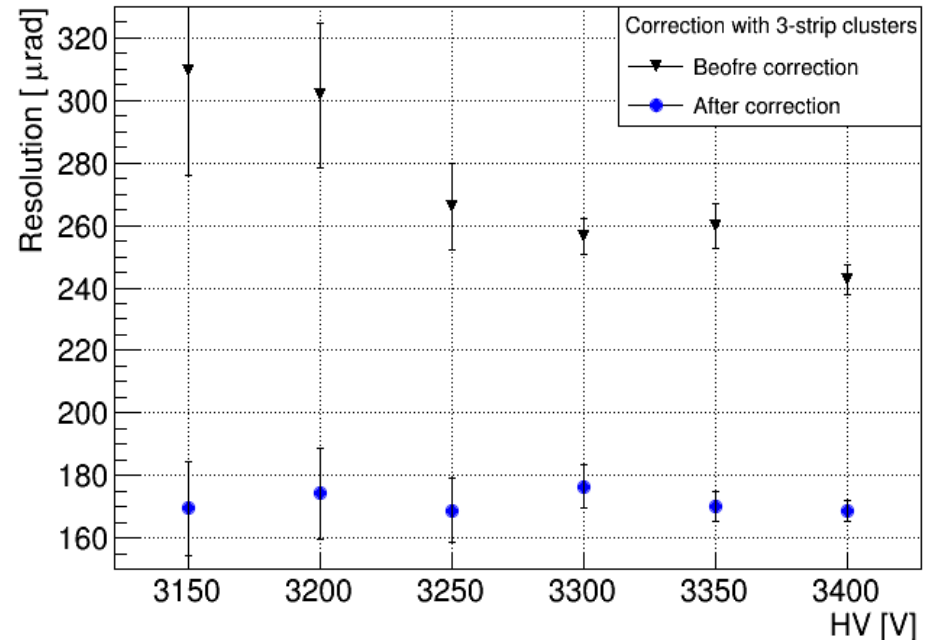


- **With 3350V data;**
- **3-strip clusters plot can be also fitted with the serpentine function; while 2-strip clusters plot can be fitted with two polynomials (degree 3).**

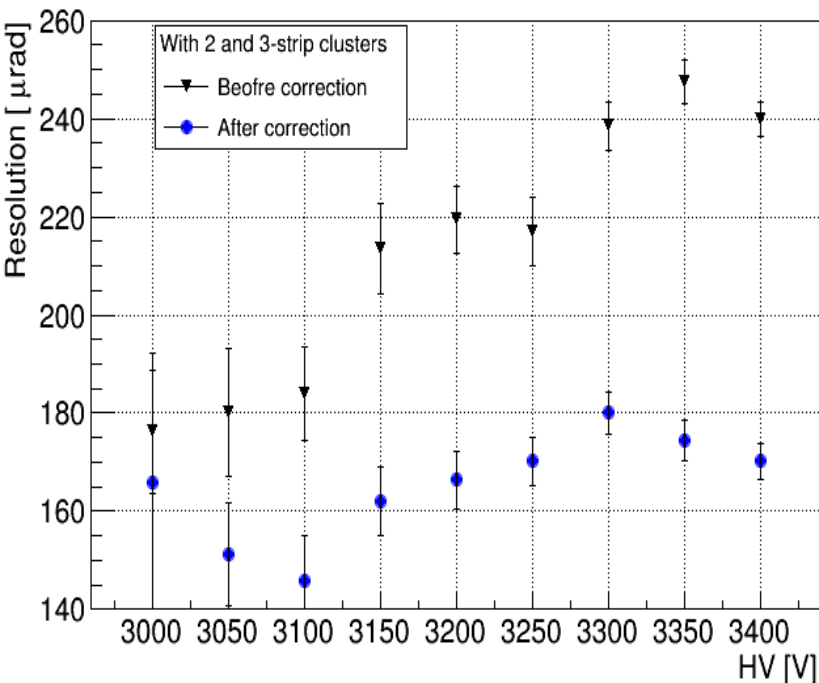
2-strip clusters



3-strip clusters



2 & 3-strip clusters



- Using the correction functions from 3350V data, calculate both exclusive and inclusive residuals. Then get the resolution from geometric mean.
- The resolutions are similar to the previous ones corrected from “HV combined data”.

Summary

- The new correction method builds 2D map of residual vs. η , with information of reference detectors. Then fits the curves with certain functions for different cluster sizes.
- The resolution results show big (& consistent for all HV points) improvement for both 2 and 3- strip clusters.
- Personal concern: the correction pulls residual towards zero on the 2D map, then in any case we'll get smaller residual width. For any probed detector in a beam test, e.g., U.Va's EIC GEM, FIT's Zigzag GEM and CMS GEM, this method improves resolution?